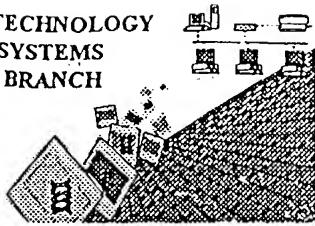


BIOTECHNOLOGY  
SYSTEMS  
BRANCH



RAW SEQUENCE LISTING  
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/650,326A  
Source: IFW3  
Date Processed by STIC: 4/26/04

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1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
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Revised 10/08/03



IFWO

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/650,326A

DATE: 04/26/2004  
TIME: 09:51:48

Input Set : A:\SEQUENCE LISTING.txt  
Output Set: N:\CRF4\04122004\J650326A.raw

3 <110> APPLICANT: CURIS INC. AND WASHINGTON UNIVERSITY  
 5 <120> TITLE OF INVENTION: CONJOINT ADMINISTRATION OF MORPHOGENS AND ACE INHIBITORS IN  
 6 TREATMENT OF CHRONIC RENAL FAILURE  
 8 <130> FILE REFERENCE: JJJ-P01-599  
 10 <140> CURRENT APPLICATION NUMBER: 10/650,326A  
 11 <141> CURRENT FILING DATE: 2003-08-28  
 13 <150> PRIOR APPLICATION NUMBER: 60/406,431  
 14 <151> PRIOR FILING DATE: 2002-08-28  
 16 <160> NUMBER OF SEQ ID NOS: 31  
 18 <170> SOFTWARE: PatentIn version 3.2  
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 21 <211> LENGTH: 139  
 22 <212> TYPE: PRT  
 23 <213> ORGANISM: Homo sapiens  
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 31 Asn Gln Glu Ala Leu Arg Met Ala Asn Val Ala Glu Asn Ser Ser Ser  
 32 20 25 30  
 35 Asp Gln Arg Gln Ala Cys Lys Lys His Glu Leu Tyr Val Ser Phe Arg  
 36 35 40 45  
 39 Asp Leu Gly Trp Gln Asp Trp Ile Ile Ala Pro Glu Gly Tyr Ala Ala  
 40 50 55 60  
 43 Tyr Tyr Cys Glu Gly Glu Cys Ala Phe Pro Leu Asn Ser Tyr Met Asn  
 44 65 70 75 80  
 47 Ala Thr Asn His Ala Ile Val Gln Thr Leu Val His Phe Ile Asn Pro  
 48 85 90 95  
 51 Glu Thr Val Pro Lys Pro Cys Cys Ala Pro Thr Gln Leu Asn Ala Ile  
 52 100 105 110  
 55 Ser Val Leu Tyr Phe Asp Asp Ser Ser Asn Val Ile Leu Lys Lys Tyr  
 56 115 120 125  
 59 Arg Asn Met Val Val Arg Ala Cys Gly Cys His  
 60 130 135  
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 65 <212> TYPE: PRT  
 66 <213> ORGANISM: Homo sapiens  
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 71 1 5 10 15  
 74 Leu Ser Ile Leu Gly Leu Pro His Arg Pro Arg Pro His Leu Gln Gly  
 75 20 25 30  
 78 Lys His Asn Ser Ala Pro Met Phe Met Leu Asp Leu Tyr Asn Ala Met

Does Not Comply  
 Corrected Diskette Needed  
 JP 6,911-14

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/650,326A

DATE: 04/26/2004  
TIME: 09:51:48

Input Set : A:\SEQUENCE LISTING.txt  
Output Set: N:\CRF4\04122004\J650326A.raw

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83	50	55	60
86	Tyr Lys Ala Val Phe Ser Thr Gln Gly Pro Pro Leu Ala Ser Leu Gln		
87	65	70	75
90	Asp Ser His Phe Leu Thr Asp Ala Asp Met Val Met Ser Phe Val Asn		80
91	95	90	
94	Leu		
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100	<213> ORGANISM: Homo sapiens		
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108	10	15	
109	Leu Trp Ala Pro Leu Phe Leu Leu Arg Ser Ala Leu Ala Asp Phe Ser		
112	20	25	30
113	Leu Asp Asn Glu Val His Ser Ser Phe Ile His Arg Arg Leu Arg Ser		
116	35	40	45
117	Gln Glu Arg Arg Glu Met Gln Arg Glu Ile Leu Ser Ile Leu Gly Leu		
120	50	55	60
121	Pro His Arg Pro Arg Pro His Leu Gln Gly Lys His Asn Ser Ala Pro		
124	65	70	75
125	Met Phe Met Leu Asp Leu Tyr Asn Ala Met Ala Val Glu Glu Gly Gly		80
128	85	90	95
129	Gly Pro Gly Gly Gln Gly Phe Ser Tyr Pro Tyr Lys Ala Val Phe Ser		
132	100	105	110
133	Thr Gln Gly Pro Pro Leu Ala Ser Leu Gln Asp Ser His Phe Leu Thr		
136	115	120	125
137	Asp Ala Asp Met Val Met Ser Phe Val Asn Leu Val Glu His Asp Lys		
140	130	135	140
141	Glu Phe Phe His Pro Arg Tyr His His Arg Glu Phe Arg Phe Asp Leu		
144	145	150	155
145	Ser Lys Ile Pro Glu Gly Glu Ala Val Thr Ala Ala Glu Phe Arg Ile		160
148	165	170	175
149	Tyr Lys Asp Tyr Ile Arg Glu Arg Phe Asp Asn Glu Thr Phe Arg Ile		
152	180	185	190
153	Ser Val Tyr Gln Val Leu Gln Glu His Leu Gly Arg Glu Ser Asp Leu		
156	195	200	205
157	Phe Leu Leu Asp Ser Arg Thr Leu Trp Ala Ser Glu Glu Gly Trp Leu		
160	210	215	220
161	Val Phe Asp Ile Thr Ala Thr Ser Asn His Trp Val Val Asn Pro Arg		
164	225	230	235
165	His Asn Leu Gly Leu Gln Leu Ser Val Glu Thr Leu Asp Gly Gln Ser		240
168	245	250	255
169	Ile Asn Pro Lys Leu Ala Gly Leu Ile Gly Arg His Gly Pro Gln Asn		
172	260	265	270
173	Lys Gln Pro Phe Met Val Ala Phe Phe Lys Ala Thr Glu Val His Phe		
	275	280	285

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/650,326A

DATE: 04/26/2004

TIME: 09:51:48

Input Set : A:\SEQUENCE LISTING.txt

Output Set: N:\CRF4\04122004\J650326A.raw

176 Arg Ser Ile Arg Ser Thr Gly Ser Lys Gln Arg Ser Gln Asn Arg Ser  
 177 290 295 300  
 180 Lys Thr Pro Lys Asn Gln Glu Ala Leu Arg Met Ala Asn Val Ala Glu  
 181 305 310 315 320  
 184 Asn Ser Ser Ser Asp Gln Arg Gln Ala Cys Lys Lys His Glu Leu Tyr  
 185 325 330 335  
 188 Val Ser Phe Arg Asp Leu Gly Trp Gln Asp Trp Ile Ile Ala Pro Glu  
 189 340 345 350  
 192 Gly Tyr Ala Ala Tyr Tyr Cys Glu Gly Glu Cys Ala Phe Pro Leu Asn  
 193 355 360 365  
 196 Ser Tyr Met Asn Ala Thr Asn His Ala Ile Val Gln Thr Leu Val His  
 197 370 375 380  
 200 Phe Ile Asn Pro Glu Thr Val Pro Lys Pro Cys Cys Ala Pro Thr Gln  
 201 385 390 395 400  
 204 Leu Asn Ala Ile Ser Val Leu Tyr Phe Asp Asp Ser Ser Asn Val Ile  
 205 405 410 415  
 208 Leu Lys Lys Tyr Arg Asn Met Val Val Arg Ala Cys Gly Cys His  
 209 420 425 430  
 212 <210> SEQ ID NO: 4  
 213 <211> LENGTH: 139  
 214 <212> TYPE: PRT  
 215 <213> ORGANISM: Mus musculus  
 217 <400> SEQUENCE: 4  
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 223 Asn Gln Glu Ala Leu Arg Met Ala Ser Val Ala Glu Asn Ser Ser Ser  
 224 20 25 30  
 227 Asp Gln Arg Gln Ala Cys Lys Lys His Glu Leu Tyr Val Ser Phe Arg  
 228 35 40 45  
 231 Asp Leu Gly Trp Gln Asp Trp Ile Ile Ala Pro Glu Gly Tyr Ala Ala  
 232 50 55 60  
 235 Tyr Tyr Cys Glu Gly Glu Cys Ala Phe Pro Leu Asn Ser Tyr Met Asn  
 236 65 70 75 80  
 239 Ala Thr Asn His Ala Ile Val Gln Thr Leu Val His Phe Ile Asn Pro  
 240 85 90 95  
 243 Asp Thr Val Pro Lys Pro Cys Cys Ala Pro Thr Gln Leu Asn Ala Ile  
 244 100 105 110  
 247 Ser Val Leu Tyr Phe Asp Asp Ser Ser Asn Val Ile Leu Lys Lys Tyr  
 248 115 120 125  
 251 Arg Asn Met Val Val Arg Ala Cys Gly Cys His  
 252 130 135  
 255 <210> SEQ ID NO: 5  
 256 <211> LENGTH: 139  
 257 <212> TYPE: PRT  
 258 <213> ORGANISM: Homo sapiens  
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 262 Ala Val Arg Pro Leu Arg Arg Arg Gln Pro Lys Lys Ser Asn Glu Leu  
 263 1 5 10 15  
 266 Pro Gln Ala Asn Arg Leu Pro Gly Ile Phe Asp Asp Val His Gly Ser

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/650,326A

DATE: 04/26/2004

TIME: 09:51:48

Input Set : A:\SEQUENCE LISTING.txt

Output Set: N:\CRF4\04122004\J650326A.raw

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270	His Gly Arg Gln Val Cys Arg Arg His Glu Leu Tyr Val Ser Phe Gln			
271	35	40	45	
274	Asp Leu Gly Trp Leu Asp Trp Val Ile Ala Pro Gln Gly Tyr Ser Ala			
275	50	55	60	
278	Tyr Tyr Cys Glu Gly Glu Cys Ser Phe Pro Leu Asp Ser Cys Met Asn			
279	65	70	75	80
282	Ala Thr Asn His Ala Ile Leu Gln Ser Leu Val His Leu Met Lys Pro			
283	85	90	95	
286	Asn Ala Val Pro Lys Ala Cys Cys Ala Pro Thr Lys Leu Ser Ala Thr			
287	100	105	110	
290	Ser Val Leu Tyr Tyr Asp Ser Ser Asn Asn Val Ile Leu Arg Lys His			
291	115	120	125	
294	Arg Asn Met Val Val Lys Ala Cys Gly Cys His			
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309	Pro His Pro Asn Lys Leu Pro Gly Ile Phe Asp Asp Gly His Gly Ser			
310	20	25	30	
313	Arg Gly Arg Glu Val Cys Arg Arg His Glu Leu Tyr Val Ser Phe Arg			
314	35	40	45	
317	Asp Leu Gly Trp Leu Asp Trp Val Ile Ala Pro Gln Gly Tyr Ser Ala			
318	50	55	60	
321	Tyr Tyr Cys Glu Gly Glu Cys Ala Phe Pro Leu Asp Ser Cys Met Asn			
322	65	70	75	80
325	Ala Thr Asn His Ala Ile Leu Gln Ser Leu Val His Leu Met Lys Pro			
326	85	90	95	
329	Asp Val Val Pro Lys Ala Cys Cys Ala Pro Thr Lys Leu Ser Ala Thr			
330	100	105	110	
333	Ser Val Leu Tyr Tyr Asp Ser Ser Asn Asn Val Ile Leu Arg Lys His			
334	115	120	125	
337	Arg Asn Met Val Val Lys Ala Cys Gly Cys His			
338	130	135		
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348	Met Arg Ala Trp Leu Leu Leu Ala Val Leu Ala Thr Phe Gln Thr			
349	1	5	10	15
352	Ile Val Arg Val Ala Ser Thr Glu Asp Ile Ser Gln Arg Phe Ile Ala			
353	20	25	30	
356	Ala Ile Ala Pro Val Ala Ala His Ile Pro Leu Ala Ser Ala Ser Gly			
357	35	40	45	

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/650,326A

DATE: 04/26/2004

TIME: 09:51:48

Input Set : A:\SEQUENCE LISTING.txt  
 Output Set: N:\CRF4\04122004\J650326A.raw

360 Ser Gly Ser Gly Arg Ser Gly Ser Arg Ser Gly Gly Ala Ser Thr Ser  
 361 50 55 60  
 364 Thr Ala Leu Ala Ala Lys Ala Phe Asn Pro Phe Ser Glu Pro Ala Ser Phe  
 365 65 70 75 80  
 368 Ser Asp Ser Asp Lys Ser His Arg Ser Lys Thr Asn Lys Lys Pro Ser  
 369 85 90 95  
 372 Lys Ser Asp Ala Asn Arg Gln Phe Asn Glu Val His Lys Pro Arg Thr  
 373 100 105 110  
 376 Asp Gln Leu Glu Asn Ser Lys Asn Met Ser Lys Gln Leu Val Asn Lys  
 377 115 120 125  
 380 Pro Asn His Asn Lys Met Ala Val Lys Glu Gln Arg Ser His His Lys  
 381 130 135 140  
 384 Lys Ser His His His Arg Ser His Gln Pro Lys Gln Ala Ser Ala Ser  
 385 145 150 155 160  
 388 Thr Glu Ser His Gln Ser Ser Ser Ile Glu Ser Ile Phe Val Glu Glu  
 389 165 170 175  
 392 Pro Thr Leu Val Leu Asp Arg Glu Val Ala Ser Ile Asn Val Pro Ala  
 393 180 185 190  
 396 Asn Ala Lys Ala Ile Ile Ala Glu Gln Gly Pro Ser Thr Tyr Ser Lys  
 397 195 200 205  
 400 Glu Ala Leu Ile Lys Asp Lys Leu Lys Pro Asp Pro Ser Thr Leu Val  
 401 210 215 220  
 404 Glu Ile Glu Lys Ser Leu Leu Ser Leu Phe Asn Met Lys Arg Pro Pro  
 405 225 230 235 240  
 408 Lys Ile Asp Arg Ser Lys Ile Ile Ile Pro Glu Pro Met Lys Lys Leu  
 409 245 250 255  
 412 Tyr Ala Glu Ile Met Gly His Glu Leu Asp Ser Val Asn Ile Pro Lys  
 413 260 265 270  
 416 Pro Gly Leu Leu Thr Lys Ser Ala Asn Thr Val Arg Ser Phe Thr His  
 417 275 280 285  
 420 Lys Asp Ser Lys Ile Asp Asp Arg Phe Pro His His Arg Phe Arg  
 421 290 295 300  
 424 Leu His Phe Asp Val Lys Ser Ile Pro Ala Asp Glu Lys Leu Lys Ala  
 425 305 310 315 320  
 428 Ala Glu Leu Gln Leu Thr Arg Asp Ala Leu Ser Gln Gln Val Val Ala  
 429 325 330 335  
 432 Ser Arg Ser Ser Ala Asn Arg Thr Arg Tyr Gln Val Leu Val Tyr Asp  
 433 340 345 350  
 436 Ile Thr Arg Val Gly Val Arg Gly Gln Arg Glu Pro Ser Tyr Leu Leu  
 437 355 360 365  
 440 Leu Asp Thr Lys Thr Val Arg Leu Asn Ser Thr Asp Thr Val Ser Leu  
 441 370 375 380  
 444 Asp Val Gln Pro Ala Val Asp Arg Trp Leu Ala Ser Pro Gln Arg Asn  
 445 385 390 395 400  
 448 Tyr Gly Leu Leu Val Glu Val Arg Thr Val Arg Ser Leu Lys Pro Ala  
 449 405 410 415  
 452 Pro His His His Val Arg Leu Arg Arg Ser Ala Asp Glu Ala His Glu  
 453 420 425 430  
 456 Arg Trp Gln His Lys Gln Pro Leu Leu Phe Thr Tyr Thr Asp Asp Gly

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<221> NAME/KEY: Variant  
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<223> OTHER INFORMATION: Xaa is Tyr, Asn or Phe  
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<222> LOCATION: (26)..(26)  
<223> OTHER INFORMATION: Xaa is Glu, His, Tyr, Asp, Gln, Ala or Ser  
<220> FEATURE:  
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<223> OTHER INFORMATION: Xaa is Ile, Met, Asn, Ala, Val, Gly or Leu  
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<220> FEATURE:  
<221> NAME/KEY: Variant  
<222> LOCATION: (71) .. (71)  
<223> OTHER INFORMATION: Xaa is Ser, Ala or Pro  
<220> FEATURE:  
<221> NAME/KEY: Variant  
<222> LOCATION: (72) .. (72)  
<223> OTHER INFORMATION: Xaa is Val, Leu Met or Ile  
<220> FEATURE:  
<221> NAME/KEY: Variant  
<222> LOCATION: (74) .. (74)

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<223> OTHER INFORMATION: Xaa is Tyr or Phe  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (75)..(75)  
<223> OTHER INFORMATION: Xaa is Phe, Tyr, Leu or His  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (76)..(76)  
<223> OTHER INFORMATION: Xaa is Asp Asn or Leu  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (77)..(77)  
<223> OTHER INFORMATION: Xaa is Asp, Glu, Asn, Arg or Ser  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (78)..(78)  
<223> OTHER INFORMATION: Xaa is Ser, Gln, Asn, Tyr or Asp  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (79)..(79)  
<223> OTHER INFORMATION: Xaa is Ser, Asn, Asp, Glu or Lys  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (80)..(80)  
<223> OTHER INFORMATION: Xaa is Asn, Thr or Lys --  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (82)..(82)  
<223> OTHER INFORMATION: Xaa is Ile, Val or Asn  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (84)..(84)  
<223> OTHER INFORMATION: Xaa is Lys or Arg  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (85)..(85)  
<223> OTHER INFORMATION: Xaa is Lys, Asn, Gln, His, Arg or Val  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (86)..(86)  
<223> OTHER INFORMATION: Xaa is Tyr, Glu or His  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (87)..(87)  
<223> OTHER INFORMATION: Xaa is Arg, Gln, Glu or Pro  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (88)..(88)  
<223> OTHER INFORMATION: Xaa is Asn, Glu, Trp or Asp  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (90)..(90)  
<223> OTHER INFORMATION: Xaa is Val, thr, Ala or Ile  
<220> FEATURE:  
<221> NAME/KEY: Variant /  
<222> LOCATION: (92)..(92)  
<223> OTHER INFORMATION: Xaa is Arg, Lys, Val, Asp, Gln or Glu

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<220> FEATURE:

<221> NAME/KEY: Variant

<222> LOCATION: (93)..(93)

<223> OTHER INFORMATION: Xaa is Ala, Gly, Glu or Ser

<220> FEATURE:

<221> NAME/KEY: Variant

<222> LOCATION: (95)..(95)

<223> OTHER INFORMATION: Xaa is Gly or Ala

<220> FEATURE:

<221> NAME/KEY: Variant

<222> LOCATION: (97)..(97)

<223> OTHER INFORMATION: Xaa is His or Arg

<400> SEQUENCE: 24

Leu Xaa Xaa Xaa Phe Xaa Xaa Xaa Gly Trp Xaa Xaa Xaa Xaa Xaa  
1 5 10 15  
Pro Xaa Xaa Xaa Xaa Ala Xaa Tyr Cys Xaa Gly Xaa Cys Xaa Pro  
20 25 30  
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Asn His Ala Xaa Xaa Xaa Xaa Xaa  
35 40 45  
Xaa Cys Cys Xaa Pro  
50 55 60  
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Leu Xaa Xaa Xaa Xaa Xaa Xaa Xaa  
65 70 75 80  
Val Xaa Leu Xaa Xaa Xaa Xaa Met Xaa Val Xaa Xaa Cys Xaa Cys  
85 90 95  
Xaa

same type of error in sequences 25-30  
Sequences 25 through 30 have Xaa's  
explained, but not "Artificial Sequence".

see p. 12

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<210> SEQ ID NO 31  
<211> LENGTH: 4  
<212> TYPE: PRT  
<213> ORGANISM: Artificial  
<220> FEATURE:  
<223> OTHER INFORMATION: :  
<400> SEQUENCE: 31  
Gly Gly Pro Pro  
1

needs explanation in 12207-12237  
section

This appeared several times in  
Sequences 28 and 29

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<220> FEATURE:

<221> NAME/KEY: VARIANT

<222> LOCATION: (46) .. (46)

<223> OTHER INFORMATION: Xaa is (Asn, Lys, Val, Thr or Gin)?

also, several locations for  
Xaa showed "Serj" as an  
amino acid represented by  
Xaa (on 12237 line)

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RAW SEQUENCE LISTING ERROR SUMMARY  
 PATENT APPLICATION: US/10/650,326A

DATE: 04/26/2004  
 TIME: 09:51:49

Input Set : A:\SEQUENCE LISTING.txt  
 Output Set: N:\CRF4\04122004\J650326A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:24; Xaa Pos. 2,3,4,6,7,8,11,12,13,14,15,16,18,19,20,21,23,26,28,30,31  
 Seq#:24; Xaa Pos. 33,34,35,36,37,38,39,40,44,45,46,47,48,49,50,51,52,53,54  
 Seq#:24; Xaa Pos. 55,56,57,58,59,60,63,65,66,67,68,69,70,71,72,74,75,76,77  
 Seq#:24; Xaa Pos. 78,79,80,82,84,85,86,87,88,89,90,92,93,95,97  
 Seq#:25; Xaa Pos. 2,3,4,5,7,8,9,11,12,13,16,17,18,19,20,21,23,24,25,26,28  
 Seq#:25; Xaa Pos. 31,33,35,36,38,39,40,41,42,43,44,45,49,50,51,52,53,54,55  
 Seq#:25; Xaa Pos. 56,57,58,59,60,61,62,63,64,65,68,70,71,72,73,74,75,76,77  
 Seq#:25; Xaa Pos. 79,80,81,82,83,84,85,87,89,90,91,92,93,95,97,98,100,102  
 Seq#:26; Xaa Pos. 2,3,4,5  
 Seq#:27; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,18,19,20,21,22,23  
 Seq#:27; Xaa Pos. 24,26,28,30,31,32,33,34,35,36,37,38,39,40,41,42,43,44,45  
 Seq#:27; Xaa Pos. 46,47,48,49,50,51,52,53,54,55,56,57,58,59,60,61,63,65,66  
 Seq#:27; Xaa Pos. 67,68,69,70,71,72,74,75,76,77,78,79,80,81,82,83,84,85,86  
 Seq#:27; Xaa Pos. 87,88,89,90,91,92,93,95,97  
 Seq#:28; Xaa Pos. 2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,23,24  
 Seq#:28; Xaa Pos. 25,26,27,28,29,31,33,35,36,37,38,39,40,41,42,43,44,45,46  
 Seq#:28; Xaa Pos. 47,48,49,50,51,52,53,54,55,56,57,58,59,60,61,62,63,64,65  
 Seq#:28; Xaa Pos. 66,68,70,71,72,73,74,75,76,77,79,80,81,82,83,84,85,86,87  
 Seq#:28; Xaa Pos. 88,89,90,91,92,93,94,95,96,97,98,100,102  
 Seq#:29; Xaa Pos. 2,3,11,16,19,23,26,35,39,41,50,52,56,57,58,60,61,65,71,73  
 Seq#:29; Xaa Pos. 75,80,82,84,89,96  
 Seq#:30; Xaa Pos. 2,3

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,  
 per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:24,25,26,27,28,29,30,31

Use of <220> Feature(NEW RULES):

error explanation

Sequence(s) are missing the <220> Feature and associated headings.  
 Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence"  
 or "Unknown". Please explain source of genetic material in <220> to <223>  
 section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32)  
 (Sec.1.823 of new Rules)

Seq#:28,29,31,24,25,26,27,30

**VERIFICATION SUMMARY**  
PATENT APPLICATION: US/10/650,326A

DATE: 04/26/2004  
TIME: 09:51:49

Input Set : A:\SEQUENCE LISTING.txt  
Output Set: N:\CRF4\04122004\J650326A.raw

L:2438 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0  
M:341 Repeated in SeqNo=24  
L:2838 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0  
M:341 Repeated in SeqNo=25  
L:2893 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0  
L:3345 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0  
M:341 Repeated in SeqNo=27  
L:3380 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:28  
L:3712 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:28  
L:3828 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:28  
L:3830 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0  
M:341 Repeated in SeqNo=28  
L:4008 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:29  
L:4010 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0  
M:341 Repeated in SeqNo=29  
L:4051 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0  
L:4060 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:31, <213>  
ORGANISM:Artificial  
L:4060 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:31, <213>  
ORGANISM:Artificial  
L:4060 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:31, Line#:4060